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piece 1, NC_000913, rpsO_truB+, config: linear, direction: +, begin: 3309677, end: 3309874

...] NC_000913.rpsC

p35 5.3 bits

p10 3.3

{-----} p35-(24)-p10 3309710 Gap 2.4 bits
{-----} p35-p10 3309710 total 6.2 bits

```
{-----} p35-(25)-p10 3309711 Gap 4.0 bits  
{-----} p35-p10 3309711 total 4.0 bits
```

A sequence alignment diagram showing a DNA template strand (top) and a reference strand (bottom). The template strand has several mutations highlighted in red: a G to T at position 3309760, a C to T at position 3309770, a C to T at position 3309780, a T to A at position 3309790, a G to A at position 3309800, a G to A at position 3309810, a T to C at position 3309820, and a T to C at position 3309830. The reference strand is identical to the template strand.

p35 4.6 bi

6.0 bit

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- NC 000813 txuB

[####> orf 41 codons

[REDACTED] off 41 codons